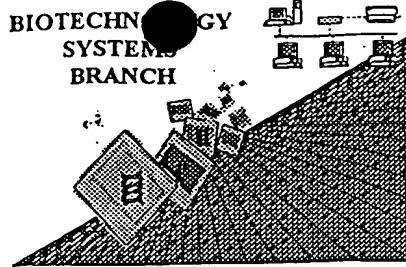
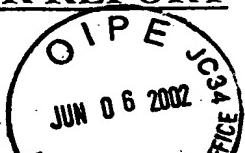


Re-run

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/647,140

Source: Patent

Date Processed by STIC: 6/18/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be downloaded from the USPTO website at the following address:
<http://www.uspto.gov/web/offices/pac/checker>

*Please mail this Raw Sequence Listing
error report with the 905.*

Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> <u>09/647,140</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTÓ SOFTWARE		
1 <input type="checkbox"/> Wrapped Nucleic Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <input type="checkbox"/> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <input type="checkbox"/> Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 <input type="checkbox"/> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <input type="checkbox"/> Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <input type="checkbox"/> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped	
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <input type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <input type="checkbox"/> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 <input type="checkbox"/> Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <input type="checkbox"/> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	

PCT09



RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/647,140

DATE: 06/18/2001
TIME: 16:26:35

Input Set : A:\FCCC Kruh ('140) Sequence Listing.txt
Output Set: N:\CRF3\06182001\I647140.raw

Does Not Comply
Corrected Diskette Needed

P.6

```

3 <110> APPLICANT: Fox Chase Cancer Center
4      Kruh, Gary D.
5      Lee, Kun
6      Belinsky, Martin G.
7      Bain, Lisa J.
9 <120> TITLE OF INVENTION: MRP-Related ABC Transporter Encoding
10     Nucleic Acids and Methods of Use Thereof
12 <130> FILE REFERENCE: FCCC 98-02
14 <140> CURRENT APPLICATION NUMBER: 09/647,140
--> 15 <141> CURRENT FILING DATE: 2001-05-21
17 <150> PRIOR APPLICATION NUMBER: PCT/US99/06644
18 <151> PRIOR FILING DATE: 1999-03-26
20 <150> PRIOR APPLICATION NUMBER: 60/079,759
21 <151> PRIOR FILING DATE: 1998-03-27
23 <150> PRIOR APPLICATION NUMBER: 60/095,153
24 <151> PRIOR FILING DATE: 1998-08-03
26 <160> NUMBER OF SEQ ID NOS: 18
28 <170> SOFTWARE: FastSEQ for Windows Version 3.0
31 <210> SEQ ID NO: 1
32 <211> LENGTH: 4231
33 <212> TYPE: DNA
34 <213> ORGANISM: Homo sapiens
36 <400> SEQUENCE: 1

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39	gcccgtgtac caggagggtga agcccaaccc gctgcaggac gcgaaacatct gtcacacgcgt	180
40	gttcttctgg tggctcaatc ccttgtttaa aattggccat aaacggagat tagaggaaga	240
41	tgtatgtat tcagtgcgtc cagaagacccg ctcacacgc ac tttggagagg agttgcaagg	300
42	gttctggat aaagaagttt taagagctga gaatgacgca cagaaggcctt cttaacaag	360
43	agcaatcata aagtgttact gaaatctta tttagtttg ggaattttta cgtaatttga	420
44	ggaaagtgcc aaagtaatcc agccatatt ttggggaaaa attattaatt attttggaaa	480
45	ttatgatccc atggattctg tggcttcaa cacagctac gcctatgcc ca ggatgtcgt	540
46	tttttcacg ctcattttgg ctatactgca tcacttat tttatcag ttcagtgtgc	600
47	tggatgagg ttacgagtag ccatgtgcca tatgattat cggaggcac ttctgttttag	660
48	taacatggcc atggggaaa caaccacagg ccagatagtc aatctgtgt ccaatgtat	720
49	gaacaagttt gatcagggtga cagtgttctt acacttcctg tggcaggac cactgcaggc	780
50	gatcgcagtg actgccctac tctggatgga gataggaata tctgcctt ctggatggc	840
51	agttctaatac attctcctgc cctgtttaaag ctgttttggg aagttgttct catcaactgag	900
52	gagtaaaact gcaactttca cggatgccag gatcaggacc atgaatgaag ttataactgg	960
53	tataaggata ataaaaatgt acgcctggga aaagtcaattt tcaaatttta ttaccaattt	1020
54	gagaaagaag gagatttcca agattctgag aagttctgc ctcaggggga tgaatttggc	1080
55	ttcggttttc agtgcaagca aaatcatctgt gtttgcgttcc accacacct acgtgtcct	1140
56	cggcaagtgtg atcacagccca gcccgtgtt cgtggcaagtg acgtgtatg gggctgtgcg	1200
57	gctgacgggtt accctcttct tcccctcagc cattgagagg gtgtcagagg caatcgtcag	1260
58	catccgaaga atccagaccc ttttgctact tgatgagata tcaacagcgca accgtcagct	1320
59	gcccgtcagat ggtaaaaaga tggtgcatgt gcaggattt actgtttttt gggataaggc	1380
60	atcagagacc ccaactctac aaggccttc cttactgtc agacactggcg aattgttagc	1440

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/647,140

DATE: 06/18/2001
TIME: 16:26:35

Input Set : A:\FCCC Kruh ('140) Sequence Listing.txt
Output Set: N:\CRF3\06182001\I647140.raw

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62	ggcccaagt	cacgggctgg tcagcgtgca	tggaagaatt gcctatgtgt	1560
63	ctgggtgttc	tcgggaactc tgaggagtaa	tatttattt gggaaagaat ataaaagga	1620
64	acgatatgaa	aaagtataa aggcttgc	tctgaaaaag gatttacagc	1680
65	tggtgatctg	actgtgatag gagatcgaaa	aaccacgctg agtggaggc	1740
66	ggtaaacctt	gcaagagcag tttatcaaga	tgctgacatc tatctcctgg	1800
67	cagtgcagta	gatgcggaaat ttagcagaca	cttggtaaa ctgtgttattt	1860
68	gcatgagaag	atcacaattt tagtgactca	tcagttgcag tacctaag	1920
69	gattctgata	ttgaaagatg gtaaaatgg	gcagaaggagg acttacactg	1980
70	atctggata	gattttggct ccctttaaa	gaaggataat gggaaatgt	2040
71	agttccagga	actcccacac taaggaatcg	taccttctca gaggcttc	2100
72	acaatcttct	agacccttct tgaaagatgg	tgctctggag agccaatgt	2160
73	cccagttaca	ctatcagagg agaaccgttc	tgaaggaaaa gttggtttc	2220
74	gaattactc	agagctgtg ctcaactggat	tgtcttcatt ttcttattt	2280
75	tgcaagtcag	gttgcctatg tgcttcaaga	ttggggctt tcatactgg	2340
76	aagtatgta	aatgtcaactg taaatggagg	aggaaatgt accgagaagc	2400
77	ctggtaactt	ggaatttattt cagtttaac	tgttagtacc gtttttttgc	2460
78	atctctattt	gtattctacg tccttggtaa	ctcttcacaa actttgcaca	2520
79	tgagtcattt	ctgaaagtc cggattttt	cttggataga aatccaaatgt	2580
80	aaatgtttc	tccaaagaca ttggacactt	ggatgatttgc gttttttaga	2640
81	tttcatccag	acattgtac aagtgggtgg	tgtggctctt gtggctgtgg	2700
82	ttggatcgca	atacccttgg ttccccttgg	aatcattttc atttttctt	2760
83	tttggaaacg	tcaagagatg tgaagegcct	ggaatctaca actcggagtc	2820
84	ccacttgtca	tcttctctcc agggctctg	gaccatcegg gatataaaag	2880
85	gtgtcaggaa	ctgtttgtatg cacaccaggaa	tttacattca gaggcttgg	2940
86	gacaacgtcc	cgctgggtcg ccgtccgtct	ggatgcattc tggccatgt	3000
87	cgttgcctt	gggtccctga ttctggcaaa	aactctggat gcccggcagg	3060
88	actgtcctat	gccttcacgc tcatggggat	gtttcaatgg tggatcgac	3120
89	agttgagaat	atgatgatct cagtagaaag	ggtcatgtt tacacagacc	3180
90	agcaccttgg	aatatcaga aaccccacc	accagcctgg cccatgt	3240
91	ctttgacaat	gtgaacttca tgcatacg	aggtggcct ctgttactgt	3300
92	agcacttcatt	aaatcacaag aaaagttgg	cattgtggaa agaaccggag	3360
93	ttccctcatc	tcagccctt ttagattgtc	agaaccggaa ggtaaaattt	3420
94	gatcttgcata	actgaaattt gacttcacga	ggattgttcaaa tcatacctca	3480
95	ggaacctgtt	ttgttcaatg gaacaatgt	aaaaacctg gatcccttta	3540
96	ggatgaggaa	ctgtggatg ctttacaaga	ggtacaactt aaagaaacca	3600
97	tcctggtaaa	atggatactg aattagcaga	atcaggatcc aatttttagt	3660
98	acaactggtg	tgccttgcca gggcaattt	caggaaaaat cagatattgt	3720
99	agcgacggca	aatgtggatc caagaactgt	tgatgttata caaaaaaaaa	3780
100	atttgccccac	tgcaccgtgc taaccattgc	acacagattt aacaccat	3840
101	caagataatg	gttttagatt cagggaaat	gttggccgt atgtttgt	3900
102	gcaaaaataaa	gagggctat ttacaagat	ggtcaacaa ctggcaagg	3960
103	tgccttcact	gaaacagcaa aacaggatata	cttcaaaaga aattatccac	4020
104	cactgaccac	atggttacaa acacttccaa	tggacagccc tgcacctaa	4080
105	gacagcactg	tgaatccaaac caaaaatgtca	agtccgttcc gaaggcat	4140
106	tttggactat	gtaaaccaca ttgtactttt	tttactttt gcaacaaata	4200
107	caagatgta	tttcatttgc atatttctcc	c	4231
110	<210>	SEQ ID NO: 2		
111	<211>	LENGTH: 1325		

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/647,140

DATE: 06/18/2001
TIME: 16:26:35

Input Set : A:\FCCC Kruh ('140) Sequence Listing.txt
Output Set: N:\CRF3\06182001\I647140.raw

112 <212> TYPE: PRT
 113 <213> ORGANISM: Homo sapiens
 115 <400> SEQUENCE: 2
 116 Met Leu Pro Val Tyr Gln Glu Val Lys Pro Asn Pro Leu Gln Asp Ala
 117 1 5 10 15
 118 Asn Ile Cys Ser Arg Val Phe Phe Trp Trp Leu Asn Pro Leu Phe Lys
 119 20 25 30
 120 Ile Gly His Lys Arg Arg Leu Glu Glu Asp Asp Met Tyr Ser Val Leu
 121 35 40 45
 122 Pro Glu Asp Arg Ser Gln His Leu Gly Glu Glu Leu Gln Gly Phe Trp
 123 50 55 60
 124 Asp Lys Glu Val Leu Arg Ala Glu Asn Asp Ala Gln Lys Pro Ser Leu
 125 65 70 75 80
 127 Thr Arg Ala Ile Ile Lys Cys Tyr Trp Lys Ser Tyr Leu Val Leu Gly
 128 85 90 95
 129 Ile Phe Thr Leu Ile Glu Glu Ser Ala Lys Val Ile Gln Pro Ile Phe
 130 100 105 110
 131 Leu Gly Lys Ile Ile Asn Tyr Phe Glu Asn Tyr Asp Pro Met Asp Ser
 132 115 120 125
 133 Val Ala Leu Asn Thr Ala Tyr Ala Tyr Ala Thr Val Leu Thr Phe Cys
 134 130 135 140
 135 Thr Leu Ile Leu Ala Ile Leu His His Leu Tyr Phe Tyr His Val Gln
 136 145 150 155 160
 137 Cys Ala Gly Met Arg Leu Arg Val Ala Met Cys His Met Ile Tyr Arg
 138 165 170 175
 139 Lys Ala Leu Arg Leu Ser Asn Met Ala Met Gly Lys Thr Thr Gly
 140 180 185 190
 141 Gln Ile Val Asn Leu Leu Ser Asn Asp Val Asn Lys Phe Asp Gln Val
 142 195 200 205
 143 Thr Val Phe Leu His Phe Leu Trp Ala Gly Pro Leu Gln Ala Ile Ala
 144 210 215 220
 145 Val Thr Ala Leu Leu Trp Met Glu Ile Gly Ile Ser Cys Leu Ala Gly
 146 225 230 235 240
 147 Met Ala Val Leu Ile Ile Leu Leu Pro Leu Gln Ser Cys Phe Gly Lys
 148 245 250 255
 149 Leu Phe Ser Ser Leu Arg Ser Lys Thr Ala Thr Phe Thr Asp Ala Arg
 150 260 265 270
 151 Ile Arg Thr Met Asn Glu Val Ile Thr Gly Ile Arg Ile Ile Lys Met
 152 275 280 285
 153 Tyr Ala Trp Glu Lys Ser Phe Ser Asn Leu Ile Thr Asn Leu Arg Lys
 154 290 295 300
 155 Lys Glu Ile Ser Lys Ile Leu Arg Ser Ser Cys Leu Arg Gly Met Asn
 156 305 310 315 320
 157 Leu Ala Ser Phe Phe Ser Ala Ser Lys Ile Ile Val Phe Val Thr Phe
 158 325 330 335
 159 Thr Thr Tyr Val Leu Leu Gly Ser Val Ile Thr Ala Ser Arg Val Phe
 160 340 345 350
 161 Val Ala Val Thr Leu Tyr Gly Ala Val Arg Leu Thr Val Thr Leu Phe
 162 355 360 365

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/647,140

DATE: 06/18/2001
TIME: 16:26:35

Input Set : A:\FCCC Kruh ('140) Sequence Listing.txt
Output Set: N:\CRF3\06182001\I647140.raw

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163 Phe Pro Ser Ala Ile Glu Arg Val Ser Glu Ala Ile Val Ser Ile Arg
164      370      375      380
165 Arg Ile Gln Thr Phe Leu Leu Leu Asp Glu Ile Ser Gln Arg Asn Arg
166      385      390      395      400
167 Gln Leu Pro Ser Asp Gly Lys Lys Met Val His Val Gln Asp Phe Thr
168      405      410      415
169 Ala Phe Trp Asp Lys Ala Ser Glu Thr Pro Thr Leu Gln Gly Leu Ser
170      420      425      430
171 Phe Thr Val Arg Pro Gly Glu Leu Leu Ala Val Val Gly Pro Val Gly
172      435      440      445
173 Ala Gly Lys Ser Ser Leu Leu Ser Ala Val Leu Gly Glu Leu Ala Pro
174      450      455      460
175 Ser His Gly Leu Val Ser Val His Gly Arg Ile Ala Tyr Val Ser Gln
176      465      470      475      480
177 Gln Pro Trp Val Phe Ser Gly Thr Leu Arg Ser Asn Ile Leu Phe Gly
178      485      490      495
179 Lys Lys Tyr Glu Lys Glu Arg Tyr Glu Lys Val Ile Lys Ala Cys Ala
180      500      505      510
181 Leu Lys Lys Asp Leu Gln Leu Leu Glu Asp Gly Asp Leu Thr Val Ile
182      515      520      525
183 Gly Asp Arg Gly Thr Pro Leu Ser Gly Gly Gln Lys Ala Arg Val Asn
184      530      535      540
185 Leu Ala Arg Ala Val Tyr Gln Asp Ala Asp Ile Tyr Leu Leu Asp Asp
186      545      550      555      560
187 Pro Leu Ser Ala Val Asp Ala Glu Val Ser Arg His Leu Phe Glu Leu
188      565      570      575
189 Cys Ile Cys Gln Ile Leu His Glu Lys Ile Thr Ile Leu Val Thr His
190      580      585      590
191 Gln Leu Gln Tyr Leu Lys Ala Ala Ser Gln Ile Leu Ile Leu Lys Asp
192      595      600      605
193 Gly Lys Met Val Gln Lys Gly Thr Tyr Thr Glu Phe Leu Lys Ser Gly
194      610      615      620
195 Ile Asp Phe Gly Ser Leu Leu Lys Lys Asp Asn Glu Glu Ser Glu Gln
196      625      630      635      640
197 Pro Pro Val Pro Gly Thr Pro Thr Leu Arg Asn Arg Thr Phe Ser Glu
198      645      650      655
199 Ser Ser Val Trp Ser Gln Gln Ser Ser Arg Pro Ser Leu Lys Asp Gly
200      660      665      670
201 Ala Leu Glu Ser Gln Asp Thr Glu Asn Val Pro Val Thr Leu Ser Glu
202      675      680      685
203 Glu Asn Arg Ser Glu Gly Lys Val Gly Phe Gln Ala Tyr Lys Asn Tyr
204      690      695      700
205 Phe Arg Ala Gly Ala His Trp Ile Val Phe Ile Phe Leu Ile Leu Leu
206      705      710      715      720
207 Asn Thr Ala Ala Gln Val Ala Tyr Val Leu Gln Asp Trp Trp Leu Ser
208      725      730      735
209 Tyr Trp Ala Asn Lys Gln Ser Met Leu Asn Val Thr Val Asn Gly Gly
210      740      745      750
211 Gly Asn Val Thr Glu Lys Leu Asp Leu Asn Trp Tyr Leu Gly Ile Tyr
212

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09/647,140

5

<210> 9
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Sequence source:/note="synthetic construct"

<400> 9

ctdgtgdgc^dg tgtdgg

see item 9 on Error Summary Sheet

18

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/647,140

DATE: 06/18/2001
TIME: 16:26:36

Input Set : A:\FCCC Kruh ('140) Sequence Listing.txt
Output Set: N:\CRF3\06182001\I647140.raw

15 M:271 C: Current Filing Date differs, Replaced Current Filing Date
1174 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:9
1174 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9
1174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
1259 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:16
1259 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:16
1259 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
1270 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:17
1270 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:17
1270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
1282 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:18
1282 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:18
1282 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18